

JPL seq list.ST25final
SEQUENCE LISTING

<110> Emtage, Peter C.R.
Tang, Y. Tom
Zhao, Q.
Liu, C.
Drmanac, R. T.

<120> MATERIALS AND METHODS RELATING TO THERAPY AND DIAGNOSIS USING TARGETING OF
CELLS THAT EXPRESS JPL POLYPEPTIDES

<130> NUV0-05

<140> NOT YET ASSIGNED

<141> 2003-07-24

<160> 12

<170> PatentIn version 3.1

<210> 1

<211> 2168

<212> DNA

<213> Homo sapiens

<400> 1
attattacag ctctgtgagg cagaggggta cctgtgaaga acctagattc cgggaatgcg 60
ccgcagccct catcgagggc tcggccacag aggtgtacgc gggcgagtgg cgcgagatc 120
ggcgagcggg cttcggcgtc agccagcgct ccaacgggct gcgctacgag ggcgagtggc 180
tgggcaaccg gcggcacggc tacggggcgca ccacccgccc cgacggctcc cgcgaggagg 240
gcaagtacaa gcgcaaccgg ctggtgcacg gcggggcgct ccgcagtctc ctgcctctgg 300
cccttcggcg gggcaagggt aaggagaagg tggacagggc tgtcgagggc gcccgtcgag 360
ccgtgagtgc tgcccgtcag cgccaggaga tcgccgctgc cagggcagca gacgccctcc 420
taaaggcagt ggcagccagc agtgtcgctg agaaggccgt ggaggcagct cgaatggcca 480
aactgatagc ccaggacctg cagcccatgc tagaggcccc aggccgcaga cccaggcagg 540
actcagaagg ttccgacacg gagcccctgg atgaggacag ccctggggta tatgagaacg 600
gactgacccc ctgagaggga tcccctgaac tgcccagcag tcctgcctcc tcccgccaac 660
cctggcgacc ccctgcctgc cggagcccac tgcctcctgg aggggaccag ggtcccttct 720
ccagcccca aagcttggcct gaggagtggg ggggggcagg cgcacaggca gaggaactag 780
ctggctatga ggctgaggat gaggctggga tgcaagggcc agggcccaga gacggttccc 840
cactcctcgg aggtgcagc gacagttcag gaagtcttcg agaggaggag ggggaggatg 900
aagagcccct gccccgctg agggccccag caggcacgga gcctgagccc atcgccatgc 960
tggtcctgag gggctcgtcc tcgaggggtc ctgatgctgg gtgcctgaca gaagagctcg 1020
gggagcccgc tgcaaccgag aggcctgccc agccgggagc tgccaacccc ctggtggtgg 1080
gagccgtggc cctcctggac ctgagcctgg cattcctgtt ctcccagctc ctcacctgag 1140

JPL seq list.ST25final

gctacttctt ggccctgggtc tggcttttggg tgcgtgcctc ttcacccctt tgacctgcct	1200
tttttctctt ctctcttcc tggctgtgtt ttctcctatc tttctttctc ttcttccttt	1260
cttttctgtg ctcttttgtt tttttctctc gctttttctt tccctgtctt ctttcagatt	1320
atctcatttc ttctggatct gtctctgtat tcctcactcc cttcccatc ccaacccctt	1380
ctttctctag attgtttaca tatgaagggc ttttctctct cagagttgct gtcttctctg	1440
agacacacaa atctaagtca gaccattgct ccacgccctc ccaccttttc tttagacctc	1500
aacttcgctg cgggtggggg tttggtgtcc taaggagact cctggaagct gaatggagag	1560
gaggaagaaa atgaagaagg agtgattgaa tgtcgggcaa ggcaactggct gagctgctgt	1620
ggctccctag cctaaggggc ctgctgtccc tctgaggcct agtgaaaaag ctgcaggagg	1680
tgcacccctc acctctaate ttggaggcta ttatcttacc tccaagcact gagctggggt	1740
actgcccatt tccatccttc cctgaaggag agaagggaag tgaaaagtag agtaactccc	1800
cagcatttcc ctctttttct cctcatcggc cagcccctcc tccagcccc tctggtgggc	1860
atgccatgcc aagagcaacg tgtaaaggaa cagagaatat ccaatgcagt caagtccac	1920
cctgcccaga ctttgccact gacttctccc acccttctgt ctccccata atagtttatt	1980
tggttggtct ggactcactt gtggcctttg attaaattcc taaggggcct gaagaagaca	2040
tttctactgc agagggttag aggcacttga gcaaggcccc cacatcccaa ctctgggagt	2100
tgtggtggga ggaggcactt ctgggggata ggaccagaca agataacagg agctcacatg	2160
gaagcaga	2168

<210> 2
 <211> 2464
 <212> DNA
 <213> Homo sapiens

<400> 2	
atgactttgt gcgtccttga agaacagcaa agaggtcaga gtggatggtg tggagtgggc	60
aagaagatga catataggat ctggggtaga agagacaacc agcccatggc tcgggacctg	120
aagcccgcca tgcctgagtc tccactcac tccatgggct cctgtgtggc ccgagcctcc	180
ccgacgagca ccacccctg ctccacggca ccagtccea tcgaccaccc aagggctgag	240
gaacgcgagc gcacggcgca ggactggcag gcagctccac ctgcagcccc ggtgcgggat	300
ccactgggtg aagccagctg ggctcctgag tctgagcctc ccctgggccc catccccga	360
aaaaccccc gaaaaacccc ccgaaaaacc gtctgcctac cttggtccga tctccccgt	420
acatattcta cccccaccc ccatcccttt gccagagacg ccgaagaacc ggggcagagg	480
ggctctgaca gcagccaggg aaacgggccc ccagcaggct ctccagtcac aggatgcgct	540
gagccgcccg ggggctgagg ccgcgccaac tacatgcatg tccccgggg gcaagtccga	600

JPL seq list.ST25final

ctttgacgac	gggggctgct	acgtgggggg	ctgggaggcg	gggcgggcac	atggctacgg	660
cgtgtgcacg	ggccccggcg	cccagggcga	gtacagcggc	tgctgggcac	acggcttcga	720
gtcactgggc	gtcttcacgg	ggccccggcg	acacagctac	cagggccact	ggcagcaggg	780
caagcgcgaa	gggctgggcg	tggagcgcga	gagccgctgg	acgtaccgcg	gcgagtggct	840
gggcgggctg	aaggggcgca	gcggcgtgtg	ggaaagcgtg	tccggcctgc	gctacgccgg	900
gctctggaag	gacggtttcc	aggacggcta	cggcactgag	acctactccg	acggaggcac	960
ctaccagggc	cagtggcagg	ccgggaagcg	ccacggctac	ggggtacgcc	agagtgtgcc	1020
ctaccatcag	gcggcgctgc	tgcgtcgcgc	ccgccgcacc	tccctggatt	ccggccacag	1080
cgaccccccg	acgccacccc	cgccccctgc	cttgccgggc	gacgagggag	gcagccccgc	1140
ctcgggctcc	cggggcggct	tcgtgctggc	cggggccggg	gacgccgacg	gcgcgtcgtc	1200
ccgaaagcgc	actccggcgg	ccggcggtatt	ctttcgccgt	tcgtgctgc	tcagcgggct	1260
ccgagcgggc	ggacgtcgca	gtccctggg	cagcaagcga	ggctccctgc	gcagcgaggt	1320
gagcagcgag	gtgggcagca	ccggaccgcc	cggctcggag	gccagcgggc	ccccggccgc	1380
agcgccgccc	gccctcatcg	agggtcggc	cacagaggtg	tacgcgggcg	agtggcgcg	1440
agatcggcgc	agcggcttcg	gcgtcagcca	gcgctccaac	gggctgcgct	acgagggcga	1500
gtggctgggc	aaccggcggc	acggctacgg	gcgcaccacc	cgccccgacg	gctcccgcga	1560
ggagggcaag	tacaagcgca	accggctggt	gcacggcggg	cgcgccgca	gtctcctgcc	1620
tctggccctt	cggcggggca	agggttaagga	gaaggtggac	agggctgtcg	agggcgcccc	1680
tcgagccgtg	agtgtgccc	gtcagcgcca	ggagatcgcc	gctgccaggg	cagcagacgc	1740
cctcctaaag	gcagtggcag	ccagcagtgt	cgctgagaag	gccgtggagg	cagctcgaat	1800
ggccaaactg	atagcccagg	acctgcagcc	catgctagag	gccccaggcc	gcagaccag	1860
gcaggactca	gaaggttccg	acacggagcc	cctggatgag	gacagccctg	gggtatatga	1920
gaacggactg	acccccctag	agggatcccc	tgaactgccc	agcagtcctg	cctcctcccc	1980
ccaaccctgg	cgacccccctg	cctgccggag	cccactgcct	cctggagggg	accagggctc	2040
cttctccagc	cccaaagctt	ggcctgagga	gtgggggggg	gcaggcgcac	aggcagagga	2100
actagctggc	tatgaggctg	aggatgaggc	tgggatgcaa	gggccagggc	ccagagacgg	2160
ttccccactc	ctcggaggct	gcagcgacag	ttcaggaagt	cttcgagagg	aggaggggga	2220
ggatgaagag	cccctgcccc	cgctgagggc	cccagcaggc	acggagcctg	agcccatcgc	2280
catgctggte	ctgaggggct	cgctcctcag	gggtcctgat	gctgggtgcc	tgacagaaga	2340
gctcggggag	cccgtgcaa	ccgagaggcc	tgcccagccg	ggagctgcca	acccccctggt	2400
ggtgggagcc	gtggccctcc	tggacctcag	cctggcattc	ctgttctccc	agctcctcac	2460

ctga

2464

<210> 3
 <211> 3065
 <212> DNA
 <213> Homo sapiens

<220>
 <221> CDS
 <222> (72)..(1955)
 <223>

<400> 3
 gccccagca ggctctccag tcccaggatg cgctgagccg ccgggggggct gaggccgcgc 60
 caactacatg c atg tcc ccc ggg ggc aag ttc gac ttt gac gac ggg ggc 110
 Met Ser Pro Gly Gly Lys Phe Asp Phe Asp Asp Gly Gly
 1 5 10
 tgc tac gtg ggg ggc tgg gag gcg ggg cgg gca cat ggc tac ggc gtg 158
 Cys Tyr Val Gly Gly Trp Glu Ala Gly Arg Ala His Gly Tyr Gly Val
 15 20 25
 tgc acg ggc ccc ggc gcc cag ggc gag tac agc ggc tgc tgg gca cac 206
 Cys Thr Gly Pro Gly Ala Gln Gly Glu Tyr Ser Gly Cys Trp Ala His
 30 35 40 45
 ggc ttc gag tca ctg ggc gtc ttc acg ggg ccc ggc gga cac agc tac 254
 Gly Phe Glu Ser Leu Gly Val Phe Thr Gly Pro Gly Gly His Ser Tyr
 50 55 60
 cag ggc cac tgg cag cag ggc aag cgc gaa ggg ctg ggc gtg gag cgc 302
 Gln Gly His Trp Gln Gln Gly Lys Arg Glu Gly Leu Gly Val Glu Arg
 65 70 75
 aag agc cgc tgg acg tac cgc ggc gag tgg ctg ggc ggg ctg aag ggc 350
 Lys Ser Arg Trp Thr Tyr Arg Gly Glu Trp Leu Gly Gly Leu Lys Gly
 80 85 90
 cgc agc ggc gtg tgg gaa agc gtg tcc ggc ctg cgc tac gcc ggg ctc 398
 Arg Ser Gly Val Trp Glu Ser Val Ser Gly Leu Arg Tyr Ala Gly Leu
 95 100 105
 tgg aag gac ggt ttc cag gac ggc tac ggc act gag acc tac tcc gac 446
 Trp Lys Asp Gly Phe Gln Asp Gly Tyr Gly Thr Glu Thr Tyr Ser Asp
 110 115 120 125
 gga ggc acc tac cag ggc cag tgg cag gcc ggg aag cgc cac ggc tac 494
 Gly Gly Thr Tyr Gln Gly Gln Trp Gln Ala Gly Lys Arg His Gly Tyr
 130 135 140
 ggg gta cgc cag agt gtg ccc tac cat cag gcg gcg ctg ctg cgc tcg 542
 Gly Val Arg Gln Ser Val Pro Tyr His Gln Ala Ala Leu Leu Arg Ser
 145 150 155
 ccc cgc cgc acc tcc ctg gat tcc ggc cac agc gac ccc ccg acg cca 590
 Pro Arg Arg Thr Ser Leu Asp Ser Gly His Ser Asp Pro Pro Thr Pro
 160 165 170
 ccc ccg ccc ctg ccc ttg ccg ggc gac gag gga ggc agc ccc gcc tcg 638
 Pro Pro Pro Leu Pro Leu Pro Gly Asp Glu Gly Gly Ser Pro Ala Ser
 175 180 185

JPL seq list.ST25final

ggc tcc cgg ggc ggc ttc gtg ctg gcc ggg ccc ggg gac gcc gac ggc Gly Ser Arg Gly Gly Phe Val Leu Ala Gly Pro Gly Asp Ala Asp Gly 190 195 200 205	686
gcg tcg tcc cga aag cgc act ccg gcg gcc ggc gga ttc ttt cgc cgt Ala Ser Ser Arg Lys Arg Thr Pro Ala Ala Gly Gly Phe Phe Arg Arg 210 215 220	734
tcg ctg ctg ctc agc ggc ctc cga gcg ggc gga cgt cgc agc tcc ctg Ser Leu Leu Leu Ser Gly Leu Arg Ala Gly Gly Arg Arg Ser Ser Leu 225 230 235	782
ggc agc aag cga ggc tcc ctg cgc agc gag gtg agc agc gag gtg ggc Gly Ser Lys Arg Gly Ser Leu Arg Ser Glu Val Ser Ser Glu Val Gly 240 245 250	830
agc acc gga ccg ccc ggc tcg gag gcc agc ggg ccc ccg gcc gca gcg Ser Thr Gly Pro Pro Gly Ser Glu Ala Ser Gly Pro Pro Ala Ala Ala 255 260 265	878
ccg ccc gcc ctc atc gag ggc tcg gcc aca gag gtg tac gcg ggc gag Pro Pro Ala Leu Ile Glu Gly Ser Ala Thr Glu Val Tyr Ala Gly Glu 270 275 280 285	926
tgg cgc gca gat cgg cgc agc ggc ttc ggc gtc agc cag cgc tcc aac Trp Arg Ala Asp Arg Arg Ser Gly Phe Gly Val Ser Gln Arg Ser Asn 290 295 300	974
ggg ctg cgc tac gag ggc gag tgg ctg ggc aac cgg cgg cac ggc tac Gly Leu Arg Tyr Glu Gly Glu Trp Leu Gly Asn Arg Arg His Gly Tyr 305 310 315	1022
ggg cgc acc acc cgc ccc gac ggc tcc cgc gag gag ggc aag tac aag Gly Arg Thr Thr Arg Pro Asp Gly Ser Arg Glu Glu Gly Lys Tyr Lys 320 325 330	1070
cgc aac cgg ctg gtg cac ggc ggg cgc gtc cgc agt ctc ctg cct ctg Arg Asn Arg Leu Val His Gly Gly Arg Val Arg Ser Leu Leu Pro Leu 335 340 345	1118
gcc ctt cgg cgg ggc aag gtt aag gag aag gtg gac agg gct gtc gag Ala Leu Arg Arg Gly Lys Val Lys Glu Lys Val Asp Arg Ala Val Glu 350 355 360 365	1166
ggc gcc cgt cga gcc gtg agt gct gcc cgt cag cgc cag gag atc gcc Gly Ala Arg Arg Ala Val Ser Ala Ala Arg Gln Arg Gln Glu Ile Ala 370 375 380	1214
gct gcc agg gca gca gac gcc ctc cta aag gca gtg gca gcc agc agt Ala Ala Arg Ala Ala Asp Ala Leu Leu Lys Ala Val Ala Ala Ser Ser 385 390 395	1262
gtc gct gag aag gcc gtg gag gca gct cga atg gcc aaa ctg ata gcc Val Ala Glu Lys Ala Val Glu Ala Ala Arg Met Ala Lys Leu Ile Ala 400 405 410	1310
cag gac ctg cag ccc atg cta gag gcc cca ggc cgc aga ccc agg cag Gln Asp Leu Gln Pro Met Leu Glu Ala Pro Gly Arg Arg Pro Arg Gln 415 420 425	1358
gac tca gaa ggt tcc gac acg gag ccc ctg gat gag gac agc cct ggg Asp Ser Glu Gly Ser Asp Thr Glu Pro Leu Asp Glu Asp Ser Pro Gly 430 435 440 445	1406

JPL seq list.ST25final

430	435	440	445	
gta tat gag aac gga ctg acc ccc tca gag gga tcc cct gaa ctg ccc Val Tyr Glu Asn Gly Leu Thr Pro Ser Glu Gly Ser Pro Glu Leu Pro	450	455	460	1454
agc agt cct gcc tcc tcc cgc caa ccc tgg cga ccc cct gcc tgc cgg Ser Ser Pro Ala Ser Ser Arg Gln Pro Trp Arg Pro Pro Ala Cys Arg	465	470	475	1502
agc cca ctg cct cct gga ggg gac cag ggt ccc ttc tcc agc ccc aaa Ser Pro Leu Pro Pro Gly Gly Asp Gln Gly Pro Phe Ser Ser Pro Lys	480	485	490	1550
gct tgg cct gag gag tgg ggg ggg gca ggc gca cag gca gag gaa cta Ala Trp Pro Glu Glu Trp Gly Gly Ala Gly Ala Gln Ala Glu Glu Leu	495	500	505	1598
gct ggc tat gag gct gag gat gag gct ggg atg caa ggg cca ggg ccc Ala Gly Tyr Glu Ala Glu Asp Glu Ala Gly Met Gln Gly Pro Gly Pro	510	515	520	1646
aga gac ggt tcc cca ctc ctc gga ggc tgc agc gac agt tca gga agt Arg Asp Gly Ser Pro Leu Leu Gly Gly Cys Ser Asp Ser Ser Gly Ser	530	535	540	1694
ctt cga gag gag gag ggg gag gat gaa gag ccc ctg ccc ccg ctg agg Leu Arg Glu Glu Glu Gly Glu Asp Glu Glu Pro Leu Pro Pro Leu Arg	545	550	555	1742
gcc cca gca ggc acg gag cct gag ccc atc gcc atg ctg gtc ctg agg Ala Pro Ala Gly Thr Glu Pro Glu Pro Ile Ala Met Leu Val Leu Arg	560	565	570	1790
ggc tcg tcc tcg agg ggt cct gat gct ggg tgc ctg aca gaa gag ctc Gly Ser Ser Ser Arg Gly Pro Asp Ala Gly Cys Leu Thr Glu Glu Leu	575	580	585	1838
ggg gag ccc gct gca acc gag agg cct gcc cag ccg gga gct gcc aac Gly Glu Pro Ala Ala Thr Glu Arg Pro Ala Gln Pro Gly Ala Ala Asn	590	595	600	1886
ccc ctg gtg gtg gga gcc gtg gcc ctc ctg gac ctc agc ctg gca ttc Pro Leu Val Val Gly Ala Val Ala Leu Leu Asp Leu Ser Leu Ala Phe	610	615	620	1934
ctg ttc tcc cag ctc ctc acc tgaggctact tcctggcctg gttctggctt Leu Phe Ser Gln Leu Leu Thr	625			1985
tggttgctg cctcttcacc cctttgacct gccttttttc tcttctcctc ttcttggtg				2045
tgttttctcc tatctttctt tctcttcttc ctttcttttc tgtgctcctt tgtttttttc				2105
tctcgctttt tctttccctg tcttctttca gattatctca tttcttctgg atctgtctct				2165
gtattcctca ctcccttccc catcccaacc ctttctttct ctagattggt tacatatgaa				2225
gggcttttct ctctcagagt tgctgtcttc tctgagacac acaaattctaa gtcagaccat				2285
tgctccacgc cctccacct tttctttaga cctcaacttc gctgcgggtg ggggtttggt				2345
gtcctaagga gactcctgga agctgaatgg agaggaggaa gaaaatgaag aaggagtgat				2405

JPL seq list.ST25final

tgaatgtcgg gcaaggcact ggctgagctg ctgtggctcc ctagcctaag gggcctgctg 2465
 tccctctgag gcctagtga aaagctgcag gaggtgcatc ctccacctct aatcttggag 2525
 gctattatct tacctccaag cactgagctg gggtactgcc caattccatc cttccctgaa 2585
 ggagagaagg gaagtga aaa gtagagtaac tccccagcat tccctcttt ttctcctcat 2645
 cggccagccc ctctccagc cccctctggg ggcattgcat gccaaagagca acgtgtaaag 2705
 gaacagagaa tatccaatgc agtcaagtcc accctgcca gactttgcca ctgacttctc 2765
 ccacccttct gtctcccca taatagttaa tttggttggg ctggactcac ttgtggcctt 2825
 tgattaaatt cctaagggg ctgaagaaga ctttctact gcagaggggt agaggcactt 2885
 gagcaaggcc cccacatccc aactctggga gttgtgggtg gaggaggcac ttctggggga 2945
 taggaccaga caagataaca ggagctcaca tggaagcaga agctgtgaca agtttagtag 3005
 tcccaaatg gggtatatcc cttccccctc gtgccgaatt cttggcctcg agggccaaat 3065

<210> 4
 <211> 771
 <212> PRT
 <213> Homo sapiens

<400> 4

Met Thr Leu Cys Val Leu Glu Glu Gln Gln Arg Gly Gln Ser Gly Trp
 1 5 10 15

Cys Gly Val Gly Lys Lys Met Thr Tyr Arg Ile Trp Gly Arg Arg Asp
 20 25 30

Asn Gln Pro Met Ala Arg Asp Leu Lys Pro Ala Met Pro Glu Ser Pro
 35 40 45

Thr His Ser Met Gly Ser Cys Val Ala Arg Ala Ser Pro Thr Ser Thr
 50 55 60

Thr Pro Cys Ser Thr Ala Pro Ser Pro Ile Asp His Pro Arg Ala Glu
 65 70 75 80

Glu Arg Glu Arg Thr Ala Gln Asp Trp Gln Ala Ala Pro Pro Ala Ala
 85 90 95

Pro Val Arg Asp Pro Leu Glu Thr Pro Lys Asn Arg Gly Arg Gly Val
 100 105 110

Leu Thr Ala Ala Arg Glu Thr Gly Pro Gln Gln Ala Leu Gln Ser Gln
 115 120 125

JPL seq list.ST25final

Asp Ala Leu Ser Arg Arg Gly Ala Glu Ala Ala Pro Thr Thr Cys Met
 130 135 140
 Ser Pro Gly Gly Lys Phe Asp Phe Asp Asp Gly Gly Cys Tyr Val Gly
 145 150 155 160
 Gly Trp Glu Ala Gly Arg Ala His Gly Tyr Gly Val Cys Thr Gly Pro
 165 170 175
 Gly Ala Gln Gly Glu Tyr Ser Gly Cys Trp Ala His Gly Phe Glu Ser
 180 185 190
 Leu Gly Val Phe Thr Gly Pro Gly Gly His Ser Tyr Gln Gly His Trp
 195 200 205
 Gln Gln Gly Lys Arg Glu Gly Leu Gly Val Glu Arg Lys Ser Arg Trp
 210 215 220
 Thr Tyr Arg Gly Glu Trp Leu Gly Gly Leu Lys Gly Arg Ser Gly Val
 225 230 235 240
 Trp Glu Ser Val Ser Gly Leu Arg Tyr Ala Gly Leu Trp Lys Asp Gly
 245 250 255
 Phe Gln Asp Gly Tyr Gly Thr Glu Thr Tyr Ser Asp Gly Gly Thr Tyr
 260 265 270
 Gln Gly Gln Trp Gln Ala Gly Lys Arg His Gly Tyr Gly Val Arg Gln
 275 280 285
 Ser Val Pro Tyr His Gln Ala Ala Leu Leu Arg Ser Pro Arg Arg Thr
 290 295 300
 Ser Leu Asp Ser Gly His Ser Asp Pro Pro Thr Pro Pro Pro Pro Leu
 305 310 315 320
 Pro Leu Pro Gly Asp Glu Gly Gly Ser Pro Ala Ser Gly Ser Arg Gly
 325 330 335
 Gly Phe Val Leu Ala Gly Pro Gly Asp Ala Asp Gly Ala Ser Ser Arg
 340 345 350
 Lys Arg Thr Pro Ala Ala Gly Gly Phe Phe Arg Arg Ser Leu Leu Leu
 355 360 365
 Ser Gly Leu Arg Ala Gly Gly Arg Arg Ser Ser Leu Gly Ser Lys Arg

JPL seq list.ST25final
380

370

375

Gly Ser Leu Arg Ser Glu Val Ser Ser Glu Val Gly Ser Thr Gly Pro
385 390 395 400

Pro Gly Ser Glu Ala Ser Gly Pro Pro Ala Ala Ala Pro Pro Ala Leu
405 410 415

Ile Glu Gly Ser Ala Thr Glu Val Tyr Ala Gly Glu Trp Arg Ala Asp
420 425 430

Arg Arg Ser Gly Phe Gly Val Ser Gln Arg Ser Asn Gly Leu Arg Tyr
435 440 445

Glu Gly Glu Trp Leu Gly Asn Arg Arg His Gly Tyr Gly Arg Thr Thr
450 455 460

Arg Pro Asp Gly Ser Arg Glu Glu Gly Lys Tyr Lys Arg Asn Arg Leu
465 470 475 480

Val His Gly Gly Arg Val Arg Ser Leu Leu Pro Leu Ala Leu Arg Arg
485 490 495

Gly Lys Val Lys Glu Lys Val Asp Arg Ala Val Glu Gly Ala Arg Arg
500 505 510

Ala Val Ser Ala Ala Arg Gln Arg Gln Glu Ile Ala Ala Arg Ala
515 520 525

Ala Asp Ala Leu Leu Lys Ala Val Ala Ala Ser Ser Val Ala Glu Lys
530 535 540

Ala Val Glu Ala Ala Arg Met Ala Lys Leu Ile Ala Gln Asp Leu Gln
545 550 555 560

Pro Met Leu Glu Ala Pro Gly Arg Arg Pro Arg Gln Asp Ser Glu Gly
565 570 575

Ser Asp Thr Glu Pro Leu Asp Glu Asp Ser Pro Gly Val Tyr Glu Asn
580 585 590

Gly Leu Thr Pro Ser Glu Gly Ser Pro Glu Leu Pro Ser Ser Pro Ala
595 600 605

Ser Ser Arg Gln Pro Trp Arg Pro Pro Ala Cys Arg Ser Pro Leu Pro
610 615 620

JPL seq list.ST25final

Pro Gly Gly Asp Gln Gly Pro Phe Ser Ser Pro Lys Ala Trp Pro Glu
625 630 635 640

Glu Trp Gly Gly Ala Gly Ala Gln Ala Glu Glu Leu Ala Gly Tyr Glu
645 650 655

Ala Glu Asp Glu Ala Gly Met Gln Gly Pro Gly Pro Arg Asp Gly Ser
660 665 670

Pro Leu Leu Gly Gly Cys Ser Asp Ser Ser Gly Ser Leu Arg Glu Glu
675 680 685

Glu Gly Glu Asp Glu Glu Pro Leu Pro Pro Leu Arg Ala Pro Ala Gly
690 695 700

Thr Glu Pro Glu Pro Ile Ala Met Leu Val Leu Arg Gly Ser Ser Ser
705 710 715 720

Arg Gly Pro Asp Ala Gly Cys Leu Thr Glu Glu Leu Gly Glu Pro Ala
725 730 735

Ala Thr Glu Arg Pro Ala Gln Pro Gly Ala Ala Asn Pro Leu Val Val
740 745 750

Gly Ala Val Ala Leu Leu Asp Leu Ser Leu Ala Phe Leu Phe Ser Gln
755 760 765

Leu Leu Thr
770

<210> 5
<211> 628
<212> PRT
<213> Homo sapiens

<400> 5

Met Ser Pro Gly Gly Lys Phe Asp Phe Asp Asp Gly Gly Cys Tyr Val
1 5 10 15

Gly Gly Trp Glu Ala Gly Arg Ala His Gly Tyr Gly Val Cys Thr Gly
20 25 30

Pro Gly Ala Gln Gly Glu Tyr Ser Gly Cys Trp Ala His Gly Phe Glu
35 40 45

Ser Leu Gly Val Phe Thr Gly Pro Gly Gly His Ser Tyr Gln Gly His
50 55 60

JPL seq list.ST25final

Trp Gln Gln Gly Lys Arg Glu Gly Leu Gly Val Glu Arg Lys Ser Arg
65 70 75 80

Trp Thr Tyr Arg Gly Glu Trp Leu Gly Gly Leu Lys Gly Arg Ser Gly
85 90 95

Val Trp Glu Ser Val Ser Gly Leu Arg Tyr Ala Gly Leu Trp Lys Asp
100 105 110

Gly Phe Gln Asp Gly Tyr Gly Thr Glu Thr Tyr Ser Asp Gly Gly Thr
115 120 125

Tyr Gln Gly Gln Trp Gln Ala Gly Lys Arg His Gly Tyr Gly Val Arg
130 135 140

Gln Ser Val Pro Tyr His Gln Ala Ala Leu Leu Arg Ser Pro Arg Arg
145 150 155 160

Thr Ser Leu Asp Ser Gly His Ser Asp Pro Pro Thr Pro Pro Pro Pro
165 170 175

Leu Pro Leu Pro Gly Asp Glu Gly Gly Ser Pro Ala Ser Gly Ser Arg
180 185 190

Gly Gly Phe Val Leu Ala Gly Pro Gly Asp Ala Asp Gly Ala Ser Ser
195 200 205

Arg Lys Arg Thr Pro Ala Ala Gly Gly Phe Phe Arg Arg Ser Leu Leu
210 215 220

Leu Ser Gly Leu Arg Ala Gly Gly Arg Arg Ser Ser Leu Gly Ser Lys
225 230 235 240

Arg Gly Ser Leu Arg Ser Glu Val Ser Ser Glu Val Gly Ser Thr Gly
245 250 255

Pro Pro Gly Ser Glu Ala Ser Gly Pro Pro Ala Ala Ala Pro Pro Ala
260 265 270

Leu Ile Glu Gly Ser Ala Thr Glu Val Tyr Ala Gly Glu Trp Arg Ala
275 280 285

Asp Arg Arg Ser Gly Phe Gly Val Ser Gln Arg Ser Asn Gly Leu Arg
290 295 300

Tyr Glu Gly Glu Trp Leu Gly Asn Arg Arg His Gly Tyr Gly Arg Thr
305 310 315 320

JPL seq list.ST25final

Thr Arg Pro Asp Gly Ser Arg Glu Glu Gly Lys Tyr Lys Arg Asn Arg
325 330 335

Leu Val His Gly Gly Arg Val Arg Ser Leu Leu Pro Leu Ala Leu Arg
340 345 350

Arg Gly Lys Val Lys Glu Lys Val Asp Arg Ala Val Glu Gly Ala Arg
355 360 365

Arg Ala Val Ser Ala Ala Arg Gln Arg Gln Glu Ile Ala Ala Ala Arg
370 375 380

Ala Ala Asp Ala Leu Leu Lys Ala Val Ala Ala Ser Ser Val Ala Glu
385 390 395 400

Lys Ala Val Glu Ala Ala Arg Met Ala Lys Leu Ile Ala Gln Asp Leu
405 410 415

Gln Pro Met Leu Glu Ala Pro Gly Arg Arg Pro Arg Gln Asp Ser Glu
420 425 430

Gly Ser Asp Thr Glu Pro Leu Asp Glu Asp Ser Pro Gly Val Tyr Glu
435 440 445

Asn Gly Leu Thr Pro Ser Glu Gly Ser Pro Glu Leu Pro Ser Ser Pro
450 455 460

Ala Ser Ser Arg Gln Pro Trp Arg Pro Pro Ala Cys Arg Ser Pro Leu
465 470 475 480

Pro Pro Gly Gly Asp Gln Gly Pro Phe Ser Ser Pro Lys Ala Trp Pro
485 490 495

Glu Glu Trp Gly Gly Ala Gly Ala Gln Ala Glu Glu Leu Ala Gly Tyr
500 505 510

Glu Ala Glu Asp Glu Ala Gly Met Gln Gly Pro Gly Pro Arg Asp Gly
515 520 525

Ser Pro Leu Leu Gly Gly Cys Ser Asp Ser Ser Gly Ser Leu Arg Glu
530 535 540

Glu Glu Gly Glu Asp Glu Glu Pro Leu Pro Pro Leu Arg Ala Pro Ala
545 550 555 560

Gly Thr Glu Pro Glu Pro Ile Ala Met Leu Val Leu Arg Gly Ser Ser
565 570 575

JPL seq list.ST25final

Ser Arg Gly Pro Asp Ala Gly Cys Leu Thr Glu Glu Leu Gly Glu Pro
580 585 590

Ala Ala Thr Glu Arg Pro Ala Gln Pro Gly Ala Ala Asn Pro Leu Val
595 600 605

Val Gly Ala Val Ala Leu Leu Asp Leu Ser Leu Ala Phe Leu Phe Ser
610 615 620

Gln Leu Leu Thr
625

<210> 6
<211> 661
<212> PRT
<213> Homo sapiens

<400> 6

Met Thr Gly Gly Arg Phe Asp Phe Asp Asp Gly Gly Thr Tyr Cys Gly
1 5 10 15

Gly Trp Glu Glu Gly Lys Ala His Gly His Gly Ile Cys Thr Gly Pro
20 25 30

Lys Gly Gln Gly Glu Tyr Ser Gly Ser Trp Ser His Gly Phe Glu Val
35 40 45

Val Gly Gly Tyr Thr Trp Pro Ser Gly Asn Thr Tyr Gln Gly Tyr Trp
50 55 60

Ala Gln Gly Lys Arg His Gly Leu Gly Val Glu Thr Lys Gly Lys Trp
65 70 75 80

Met Tyr Arg Gly Glu Trp Ser His Gly Phe Lys Gly Arg Tyr Gly Val
85 90 95

Arg Gln Ser Leu Cys Thr Pro Ala Arg Tyr Glu Gly Thr Trp Ser Asn
100 105 110

Gly Leu Gln Asp Gly Tyr Gly Val Glu Thr Tyr Gly Asp Gly Gly Thr
115 120 125

Tyr Gln Gly Gln Trp Ala Gly Gly Met Arg His Gly Tyr Gly Val Arg
130 135 140

Gln Ser Val Pro Tyr Gly Met Ala Thr Val Ile Arg Ser Pro Leu Arg
145 150 155 160

JPL seq list.ST25final

Thr Ser Leu Ala Ser Leu Arg Ser Glu Gln Ser Asn Gly Ser Val Leu
165 170 175

His Asp Ala Ala Ala Ala Ala Asp Ser Pro Ala Gly Thr Arg Gly Gly
180 185 190

Phe Val Leu Asn Phe His Ala Asp Ala Glu Leu Ala Gly Lys Lys Lys
195 200 205

Gly Gly Leu Phe Arg Arg Gly Ser Leu Leu Gly Ser Met Lys Leu Arg
210 215 220

Lys Ser Glu Ser Lys Ser Ser Ile Ser Ser Lys Arg Ser Ser Val Arg
225 230 235 240

Ser Asp Ala Ala Met Ser Arg Ile Ser Ser Ser Asp Ala Asn Ser Thr
245 250 255

Ile Ser Phe Gly Asp Val Asp Cys Asp Phe Cys Pro Val Glu Asp His
260 265 270

Val Asp Ala Thr Thr Thr Glu Thr Tyr Met Gly Glu Trp Lys Asn Asp
275 280 285

Lys Arg Asn Gly Phe Gly Val Ser Glu Arg Ser Asn Gly Met Lys Tyr
290 295 300

Glu Gly Glu Trp Ala Asn Asn Lys Arg His Gly Tyr Gly Cys Thr Val
305 310 315 320

Phe Pro Asp Gly Ser Lys Glu Glu Gly Lys Tyr Lys Asn Asn Ile Leu
325 330 335

Val Arg Gly Ile Arg Lys Gln Leu Ile Pro Ile Arg His Thr Lys Thr
340 345 350

Arg Glu Lys Val Asp Arg Ala Ile Glu Gly Ala Gln Arg Ala Ala Ala
355 360 365

Met Ala Arg Thr Lys Val Glu Ile Ala Asn Ser Arg Thr Ala His Ala
370 375 380

Arg Ala Lys Ala Asp Ala Ala Asp Gln Ala Ala Leu Ala Ala Arg Gln
385 390 395 400

Glu Cys Asp Ile Ala Arg Ala Val Ala Arg Glu Leu Ser Pro Asp Phe

JPL seq list.ST25final
410

405

415

Tyr Gln Pro Gly Pro Asp Tyr Val Lys Gln Arg Phe Gln Glu Gly Val
420 425 430

Asp Ala Lys Glu Asn Pro Glu Glu Lys Val Pro Glu Lys Pro Pro Thr
435 440 445

Pro Lys Glu Ser Pro His Phe Tyr Arg Lys Gly Thr Thr Pro Pro Arg
450 455 460

Ser Pro Glu Ala Ser Pro Lys His Ser His Ser Pro Ala Ser Ser Pro
465 470 475 480

Lys Pro Leu Lys Lys Gln Asn Pro Ser Ser Gly Ala Arg Leu Asn Gln
485 490 495

Asp Lys Arg Ser Val Ala Asp Glu Gln Val Thr Ala Ile Val Asn Lys
500 505 510

Pro Leu Met Ser Lys Ala Pro Thr Lys Glu Ala Gly Ala Val Val Pro
515 520 525

Gln Ser Lys Tyr Ser Gly Arg His His Ile Pro Asn Pro Ser Asn Gly
530 535 540

Glu Leu His Ser Gln Tyr His Gly Tyr Tyr Val Lys Leu Asn Ala Pro
545 550 555 560

Gln His Pro Pro Val Asp Val Glu Asp Gly Asp Gly Ser Ser Gln Ser
565 570 575

Ser Ser Ala Leu Val His Lys Pro Ser Ala Asn Lys Trp Ser Pro Ser
580 585 590

Lys Ser Val Thr Lys Pro Val Ala Lys Glu Ser Lys Ala Glu Pro Lys
595 600 605

Ala Lys Lys Ser Glu Leu Ala Ile Pro Lys Asn Pro Ala Ser Asn Asp
610 615 620

Ser Cys Pro Ala Leu Glu Lys Glu Ala Asn Ser Gly Pro Asn Ser Ile
625 630 635 640

Met Ile Val Leu Val Met Leu Leu Asn Ile Gly Leu Ala Ile Leu Phe
645 650 655

JPL seq list.ST25final

val His Phe Leu Thr
660

<210> 7
<211> 696
<212> PRT
<213> Homo sapiens
<400> 7

Met Ser Gly Gly Arg Phe Asp Phe Asp Asp Gly Gly Ala Tyr Cys Gly
1 5 10 15

Gly Trp Glu Gly Gly Lys Ala His Gly His Gly Leu Cys Thr Gly Pro
20 25 30

Lys Gly Gln Gly Glu Tyr Ser Gly Ser Trp Asn Phe Gly Phe Glu Val
35 40 45

Ala Gly Val Tyr Thr Trp Pro Ser Gly Asn Thr Phe Glu Gly Tyr Trp
50 55 60

Ser Gln Gly Lys Arg His Gly Leu Gly Ile Glu Thr Lys Gly Arg Trp
65 70 75 80

Leu Tyr Lys Gly Glu Trp Thr His Gly Phe Lys Gly Arg Tyr Gly Ile
85 90 95

Arg Gln Ser Ser Ser Ser Gly Ala Lys Tyr Glu Gly Thr Trp Asn Asn
100 105 110

Gly Leu Gln Asp Gly Tyr Gly Thr Glu Thr Tyr Ala Asp Gly Gly Thr
115 120 125

Tyr Gln Gly Gln Phe Thr Asn Gly Met Arg His Gly Tyr Gly Val Arg
130 135 140

Gln Ser Val Pro Tyr Gly Met Ala Val Val Val Arg Ser Pro Leu Arg
145 150 155 160

Thr Ser Leu Ser Ser Leu Arg Ser Glu His Ser Asn Gly Thr Val Ala
165 170 175

Pro Asp Ser Pro Ala Ser Pro Ala Ser Asp Gly Pro Ala Leu Pro Ser
180 185 190

Pro Ala Ile Pro Arg Gly Gly Phe Ala Leu Ser Leu Leu Ala Asn Ala
195 200 205

JPL seq list.ST25final

Glu Ala Ala Ala Arg Ala Pro Lys Gly Gly Gly Leu Phe Gln Arg Gly
210 215 220

Ala Leu Leu Gly Lys Leu Arg Arg Ala Glu Ser Arg Thr Ser Val Gly
225 230 235 240

Ser Gln Arg Ser Arg Val Ser Phe Leu Lys Ser Asp Leu Ser Ser Gly
245 250 255

Ala Ser Asp Ala Ala Ser Thr Ala Ser Leu Gly Glu Ala Ala Glu Gly
260 265 270

Ala Asp Glu Ala Ala Pro Phe Glu Ala Asp Ile Asp Ala Thr Thr Thr
275 280 285

Glu Thr Tyr Met Gly Glu Trp Lys Asn Asp Lys Arg Ser Gly Phe Gly
290 295 300

Val Ser Glu Arg Ser Ser Gly Leu Arg Tyr Glu Gly Glu Trp Leu Asp
305 310 315 320

Asn Leu Arg His Gly Tyr Gly Cys Thr Thr Leu Pro Asp Gly His Arg
325 330 335

Glu Glu Gly Lys Tyr Arg His Asn Val Leu Val Lys Asp Thr Lys Arg
340 345 350

Arg Met Leu Gln Leu Lys Ser Asn Lys Val Arg Gln Lys Val Glu His
355 360 365

Ser Val Glu Gly Ala Gln Arg Ala Ala Ala Ile Ala Arg Gln Lys Ala
370 375 380

Glu Ile Ala Ala Ser Arg Thr Ser His Ala Lys Ala Lys Ala Glu Ala
385 390 395 400

Ala Glu Gln Ala Ala Leu Ala Ala Asn Gln Glu Ser Asn Ile Ala Arg
405 410 415

Thr Leu Ala Arg Glu Leu Ala Pro Asp Phe Tyr Gln Pro Gly Pro Glu
420 425 430

Tyr Gln Lys Arg Arg Leu Leu Gln Glu Ile Leu Glu Asn Ser Glu Ser
435 440 445

Leu Leu Glu Pro Pro Asp Arg Gly Ala Gly Ala Ala Gly Leu Pro Gln
450 455 460

JPL seq list.ST25final

Pro Pro Arg Glu Ser Pro Gln Leu His Glu Arg Glu Thr Pro Arg Pro
465 470 475 480

Glu Gly Gly Ser Pro Ser Pro Ala Gly Thr Pro Pro Gln Pro Lys Arg
485 490 495

Pro Arg Pro Gly Val Ser Lys Asp Gly Leu Leu Ser Pro Gly Ala Trp
500 505 510

Asn Gly Glu Pro Ser Gly Glu Gly Ser Arg Ser Val Thr Pro Ser Glu
515 520 525

Gly Ala Gly Arg Arg Ser Pro Ala Arg Pro Ala Thr Glu Arg Met Ala
530 535 540

Ile Glu Ala Leu Gln Ala Pro Pro Ala Pro Ser Arg Glu Pro Glu Val
545 550 555 560

Ala Leu Tyr Gln Gly Tyr His Ser Tyr Ala Val Arg Thr Thr Pro Pro
565 570 575

Glu Pro Pro Pro Phe Glu Asp Gln Pro Glu Pro Glu Val Ser Gly Ser
580 585 590

Glu Ser Ala Pro Ser Ser Pro Ala Thr Ala Pro Leu Gln Ala Pro Thr
595 600 605

Leu Arg Gly Pro Glu Pro Ala Arg Glu Thr Pro Ala Lys Leu Glu Pro
610 615 620

Lys Pro Ile Ile Pro Lys Ala Glu Pro Arg Ala Lys Ala Arg Lys Thr
625 630 635 640

Glu Ala Arg Gly Leu Thr Lys Ala Gly Ala Lys Lys Lys Ala Arg Lys
645 650 655

Glu Ala Ala Leu Ala Ala Glu Ala Glu Val Glu Val Glu Glu Val Pro
660 665 670

Asn Thr Ile Leu Ile Cys Met Val Ile Leu Leu Asn Ile Gly Leu Ala
675 680 685

Ile Leu Phe Val His Leu Leu Thr
690 695

<210> 8
<211> 748

JPL seq list.ST25final

<212> PRT

<213> Homo sapiens

<400> 8

Met Ser Ser Gly Gly Arg Phe Asn Phe Asp Asp Gly Gly Ser Tyr Cys
1 5 10 15

Gly Gly Trp Glu Asp Gly Lys Ala His Gly His Gly Val Cys Thr Gly
20 25 30

Pro Lys Gly Gln Gly Glu Tyr Thr Gly Ser Trp Ser His Gly Phe Glu
35 40 45

Val Leu Gly Val Tyr Thr Trp Pro Ser Gly Asn Thr Tyr Gln Gly Thr
50 55 60

Trp Ala Gln Gly Lys Arg His Gly Ile Gly Leu Glu Ser Lys Gly Lys
65 70 75 80

Trp Val Tyr Lys Gly Glu Trp Thr His Gly Phe Lys Gly Arg Tyr Gly
85 90 95

Val Arg Glu Cys Ala Gly Asn Gly Ala Lys Tyr Glu Gly Thr Trp Ser
100 105 110

Asn Gly Leu Gln Asp Gly Tyr Gly Thr Glu Thr Tyr Ser Asp Gly Gly
115 120 125

Thr Tyr Gln Gly Gln Trp Val Gly Gly Met Arg Gln Gly Tyr Gly Val
130 135 140

Arg Gln Ser Val Pro Tyr Gly Met Ala Ala Val Ile Arg Ser Pro Leu
145 150 155 160

Arg Thr Ser Ile Asn Ser Leu Arg Ser Glu His Thr Asn Gly Thr Pro
165 170 175

Leu His Pro Asp Ala Ser Pro Ala Val Asp Gly Ser Pro Ala Val Ser
180 185 190

Arg Gly Gly Phe Val Leu Val Ala His Ser Asp Ser Glu Ile Leu Lys
195 200 205

Ser Lys Lys Lys Gly Leu Phe Arg Arg Ser Leu Leu Ser Gly Leu Lys
210 215 220

Leu Arg Lys Ser Glu Ser Lys Ser Ser Leu Ala Ser Gln Arg Ser Lys
225 230 235 240

JPL seq list.ST25final

Gln Ser Ser Phe Arg Ser Glu Ala Gly Met Ser Thr Val Ser Ser Thr
245 250 255

Ala Ser Asp Ile His Ser Thr Ile Ser Leu Gly Glu Ala Glu Ala Glu
260 265 270

Leu Ala Val Ile Glu Asp Asp Ile Asp Ala Thr Thr Thr Glu Thr Tyr
275 280 285

Val Gly Glu Trp Lys Asn Asp Lys Arg Ser Gly Phe Gly Val Ser Gln
290 295 300

Arg Ser Asp Gly Leu Lys Tyr Glu Gly Glu Trp Ala Ser Asn Arg Arg
305 310 315 320

His Gly Tyr Gly Cys Met Thr Phe Pro Asp Gly Thr Lys Glu Glu Gly
325 330 335

Lys Tyr Lys Gln Asn Ile Leu Val Gly Gly Lys Arg Lys Asn Leu Ile
340 345 350

Pro Leu Arg Ala Ser Lys Ile Arg Glu Lys Val Asp Arg Ala Val Glu
355 360 365

Ala Ala Glu Arg Ala Ala Thr Ile Ala Lys Gln Lys Ala Glu Ile Ala
370 375 380

Ala Ser Arg Thr Ser His Ser Arg Ala Lys Ala Glu Ala Ala Leu Thr
385 390 395 400

Ala Ala Gln Lys Ala Gln Glu Glu Ala Arg Ile Ala Arg Ile Thr Ala
405 410 415

Lys Glu Phe Ser Pro Ser Phe Gln His Arg Glu Asn Gly Leu Glu Tyr
420 425 430

Gln Arg Pro Lys Arg Gln Thr Ser Cys Asp Asp Ile Glu Val Leu Ser
435 440 445

Thr Gly Thr Pro Leu Gln Gln Glu Ser Pro Glu Leu Tyr Arg Lys Gly
450 455 460

Thr Thr Pro Ser Asp Leu Thr Pro Asp Asp Ser Pro Leu Gln Ser Phe
465 470 475 480

Pro Thr Ser Pro Ala Ala Thr Pro Pro Pro Ala Pro Ala Ala Arg Asn

JPL seq list.ST25final

485

490

495

Lys Val Ala His Phe Ser Arg Gln Val Ser Val Asp Glu Glu Arg Gly
500 505 510

Gly Asp Ile Gln Met Leu Leu Glu Gly Arg Ala Gly Asp Cys Ala Arg
515 520 525

Ser Ser Trp Gly Glu Glu Gln Ala Gly Gly Ser Arg Gly Val Arg Ser
530 535 540

Gly Ala Leu Arg Gly Gly Leu Leu Val Asp Asp Phe Arg Thr Arg Gly
545 550 555 560

Ser Gly Arg Lys Gln Pro Gly Asn Pro Lys Pro Arg Glu Arg Arg Thr
565 570 575

Glu Ser Pro Pro Val Phe Thr Trp Thr Ser His His Arg Ala Ser Asn
580 585 590

His Ser Pro Gly Gly Ser Arg Leu Leu Glu Leu Gln Glu Glu Lys Leu
595 600 605

Ser Asn Tyr Arg Met Glu Met Lys Pro Leu Leu Arg Met Glu Thr His
610 615 620

Pro Gln Lys Arg Arg Tyr Ser Lys Gly Gly Ala Cys Arg Gly Leu Gly
625 630 635 640

Asp Asp His Arg Pro Glu Asp Arg Gly Phe Gly Val Gln Arg Leu Arg
645 650 655

Ser Lys Ala Gln Asn Lys Glu Asn Phe Arg Pro Ala Ser Ser Ala Glu
660 665 670

Pro Ala Val Gln Lys Leu Ala Ser Leu Arg Leu Gly Gly Ala Glu Pro
675 680 685

Arg Leu Leu Arg Trp Asp Leu Thr Phe Ser Pro Pro Gln Lys Ser Leu
690 695 700

Pro Val Ala Leu Glu Ser Asp Glu Glu Asn Gly Asp Glu Leu Lys Ser
705 710 715 720

Ser Thr Gly Ser Ala Pro Ile Leu Val Val Met Val Ile Leu Leu Asn
725 730 735

JPL seq list.ST25final

Ile Gly Val Ala Ile Leu Phe Ile Asn Phe Phe Ile
740 745

<210> 9
<211> 17
<212> PRT
<213> Homo sapiens

<400> 9

Leu Val Val Gly Ala Val Ala Leu Leu Asp Leu Ser Leu Ala Phe Leu
1 5 10 15

Phe

<210> 10
<211> 35
<212> PRT
<213> Homo sapiens

<400> 10

Arg Arg Thr Ser Leu Asp Ser Gly His Ser Asp Pro Pro Thr Pro Pro
1 5 10 15

Pro Pro Leu Pro Leu Pro Gly Asp Glu Gly Gly Ser Pro Ala Ser Gly
20 25 30

Ser Arg Gly
35

<210> 11
<211> 19
<212> DNA
<213> Homo sapiens

<400> 11

gcagacgccc tcctaaagg

19

<210> 12
<211> 21
<212> DNA
<213> Homo sapiens

<400> 12

gctatcagtt tggccattcg a

21